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<110> Chermesh, Chen
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Rotman, Galit
Sella-Tavor, Osnat

<120> SIM2 POLYPEPTIDES AND POLYNUCLEOTIDES ENCODING SAME AND USES
THEREOF IN DIAGNOSIS AND TREATMENT OF OVARIAN, BREAST AND LUNG
CANCERS

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<210> 8
 <211> 2823
 <212> DNA
 <213> Homo sapiens

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agcttgccaa gctgctcccg ctgccgtcgg ccatcacttc gcagctggac aaagcgtcca	180
tcatccgcct caccacgagc tacctgaaga tgcgcgccgt cttccccgaa ggtttaggag	240
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cgcacttgct gcagactttg gatggatttg tttttgtggt agcatctgat ggcaaaatca	360
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cccaccagcc gctgcaccac cacctgctcc aagagtatga gatagagagg tcgttctttc	540
ttcgaatgaa atgtgtcttg gcgaaaagga acgcgggcct gacctgcagc ggatacaagg	600
tcatccactg cagtggctac ttgaagatca ggcagtatat gctggacatg tccctgtacg	660
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cggccaagtt cgggcagccc caaggatccc cttgtgaggt ggcacgcttt ttcctgagca 1500
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 gtgggcggat cacctgaggt caggagtttg cgacaagcct gccaacaagc tgaaacccca 2640
 tctccactaa aaatacaaaa attagttggg catggtggtg agcacctgta atcccagcta 2700
 ctctggaggc tgagatagga ggatcacttg aaccggggag gtggaggttg cagtgagcta 2760
 agatcacatc actgcactcc agcctgggta acagagtgag actgtctcaa aaaaaaaaaa 2820
 aaa 2823

<210> 9
 <211> 99
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

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 atggcaaaat catgtatata tccgagaccg cttctgtcc 99

<210> 10
 <211> 19
 <212> DNA
 <213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 10

ggagctggga tcgcacttg

19

<210> 11

<211> 21

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 11

ggacagaagc ggtctcgat a

21

<210> 12

<211> 23

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 12

tttacaacat aaagcgcgatg gtg

23

<210> 13

<211> 22

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 13

gtggctactt gaagatcagg ca

22

<210> 14

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 14

cctagcagct cgtctccagc

20

<210> 15

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 15

ggtgtcctcg ccgaacct

18

<210> 16
 <211> 21
 <212> DNA
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<220>
 <223> Single strand DNA oligonucleotide

<400> 16
 tagggacctg ccctgtacat g 21

<210> 17
 <211> 17
 <212> DNA
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<220>
 <223> Single strand DNA oligonucleotide

<400> 17
 gctggcggtg tctccgc 17

<210> 18
 <211> 120
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

<400> 18
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 agcctggtgc caagctacga agcgcccgcc gccgccgtgc gcagggttcgg cgaggacacc 120

<210> 19
 <211> 101
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

<400> 19
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 gtggcaggca agcagatgtc ctctgcggag ataccgccag c 101

<210> 20
 <211> 101
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

<400> 20
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 ctaagtgtctg gggaacttca atgaaaacct ggctggtaaa g 101

<210> 21
 <211> 20
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide

<400> 21
 ctggcaagca gctggaagat 20

<210> 22
 <211> 23
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide

<400> 22
 tttcttagca ccaccacgaa gtc 23

<210> 23
 <211> 101
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

<400> 23
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 ctcttcatct tgtgttgaga cttcgtggtg gtgctaagaa a 101

<210> 24
 <211> 25
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide

<400> 24
 cagtgattat aggctttcgc tctaa 25

<210> 25
 <211> 23
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide

<400> 25
 cagggtatt ggttgaatga gta 23

<210> 26
 <211> 134
 <212> DNA
 <213> Artificial sequence

<220>

<223> Real time PCR amplicon

<400> 26

cagtgtattat aggcctttcgc tctaagatta aaaatgccct agcccacttc ttaccacaag 60

gcacacctac accccttata cccatactag ttattatcga aaccatcagc ctactcattc 120

aaccaatagc cctg 134

<210> 27

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 27

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<210> 28

<211> 20

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 28

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<210> 29

<211> 151

<212> DNA

<213> Artificial sequence

<220>

<223> Real time PCR amplicon

<400> 29

gtaacccggt gaacccatt cgtgatgggg atcggggatt gcaattattc cccatgaacg 60

aggaattccc agtaagtgcg ggtcataagc ttgcgttgat taagtcctcg ccctttgtac 120

acaccgcccg tcgtactac cgattggatg g 151

<210> 30

<211> 19

<212> DNA

<213> Artificial sequence

<220>

<223> Single strand DNA oligonucleotide

<400> 30

tgagagtgat tcgcgtggg 19

<210> 31

<211> 21

<212> DNA

<213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide
 <400> 31
 ccagggtacg aggctttcaa t 21
 <210> 32
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 <400> 33
 tgacactggc aaaacaatgc a 21
 <210> 34
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 <223> Single strand DNA oligonucleotide
 <400> 34
 ggtccttttc accagcaagc t 21
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 <223> Real time PCR amplicon
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 ggtcaaggtc gcaagcttgc tggtgaaaag gacc 94
 <210> 36
 <211> 20
 <212> DNA
 <213> Artificial sequence
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 <223> Single strand DNA oligonucleotide

<400> 36
 tgggaacaag agggcatctg

20

<210> 37
 <211> 22
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Single strand DNA oligonucleotide

<400> 37
 ccaccactgc atcaaattca tg

22

<210> 38
 <211> 86
 <212> DNA
 <213> Artificial sequence

<220>
 <223> Real time PCR amplicon

<400> 38
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60

ggatcatgaa tttgatgcag tgggtgg

86

<210> 39
 <211> 160
 <212> PRT
 <213> Homo sapiens

<400> 39

Leu Asp Gly Phe Val Phe Val Val Ala Ser Asp Gly Lys Ile Met Tyr
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Ile Ser Glu Thr Ala Ser Val His Leu Gly Leu Ser Gln Val Glu Leu
 20 25 30

Thr Gly Asn Ser Ile Tyr Glu Tyr Ile His Pro Ser Asp His Asp Glu
 35 40 45

Met Thr Ala Val Leu Thr Ala His Gln Pro Leu His His His Leu Leu
 50 55 60

Gln Glu Tyr Glu Ile Glu Arg Ser Phe Phe Leu Arg Met Lys Cys Val
 65 70 75 80

Leu Ala Lys Arg Asn Ala Gly Leu Thr Cys Ser Gly Tyr Lys Val Ile
 85 90 95

His Cys Ser Gly Tyr Leu Lys Ile Arg Gln Tyr Met Leu Asp Met Ser
 100 105 110

Leu Tyr Asp Ser Cys Tyr Gln Ile Val Gly Leu Val Ala Val Gly Gln

115 120 125
 Ser Leu Pro Pro Ser Ala Ile Thr Glu Ile Lys Leu Tyr Ser Asn Met
 130 135 140

 Phe Met Phe Arg Ala Ser Leu Asp Leu Lys Leu Ile Phe Leu Asp Ser
 145 150 155 160

 <210> 40
 <211> 178
 <212> PRT
 <213> Homo sapiens

 <400> 40
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 1 5 10 15

 Asn Gly Glu Phe Tyr Glu Leu Ala Lys Leu Leu Pro Leu Pro Ser Ala
 20 25 30

 Ile Thr Ser Gln Leu Asp Lys Ala Ser Ile Ile Arg Leu Thr Thr Ser
 35 40 45

 Tyr Leu Lys Met Arg Ala Val Phe Pro Glu Gly Glu Ala Ser Gly Gly
 50 55 60

 Arg Pro Gly Thr Leu Gly Ser Pro Ala Ala Pro Ala Gln Ala Gly Ser
 65 70 75 80

 Ala Ser Gln Pro Ala Gln Arg Gly Cys Arg Gly Leu Ala Ser Arg Ala
 85 90 95

 Gly Ala Ser Glu Gly Gly Cys Val Arg Val Phe Gly Phe Gly Ala Gly
 100 105 110

 Leu Gly Arg Gly Ala Arg Ala Leu Ala Ala Gln Ala Thr Lys Pro Ser
 115 120 125

 Pro Gly Pro Gly Leu Gly Glu Gly Glu Leu Arg Ile Val Pro Gly Ala
 130 135 140

 Gly Ser Pro Pro Ala Arg Thr Ala Ser Glu Arg Cys Glu Ser Ala Gly
 145 150 155 160

 Ile Thr Val Arg Pro Lys His Cys Arg Leu Arg Pro Gln Ser Glu His
 165 170 175

 Leu Cys

<210> 41

<211> 584
 <212> PRT
 <213> Homo sapiens

<400> 41

Met Arg Arg Trp Leu Pro Phe Pro Pro Phe Leu Pro Gly Gly Pro Ser
 1 5 10 15

Asn Leu Ala Ile Pro Asp Phe Gln Gln His Ser Val Pro Thr Gly Pro
 20 25 30

Ala Val Leu Cys Arg Gly Ala Gly Ser Ser Pro Pro Pro Ser Arg Asn
 35 40 45

Leu Thr Phe Leu Thr Pro Thr Pro Pro Ala Gly Asn His Arg Leu Gly
 50 55 60

Trp Gly Pro Leu Val Leu Gln Glu Ser Ser Ala Asp Phe Ser Pro Ser
 65 70 75 80

Leu Pro Ala Met Gly Phe Ala Arg Ser Gln Gly Pro Glu Ala Thr Leu
 85 90 95

Thr Val Asn Thr Ala Gln Gly Lys Gly Gly Ser Arg Thr Thr Ala Gln
 100 105 110

Pro Gln Glu Arg Pro Ser Ser Arg Glu Lys Asp Ser Cys Gln Gly Ala
 115 120 125

Phe Val Pro Arg Pro Ser Tyr Pro Ala Leu Pro Gly Gly Ser Cys Leu
 130 135 140

Trp Phe Ser Ala Arg Gly Thr Thr Ala Leu Arg Thr Ala Val Gly Gly
 145 150 155 160

Trp Gly Trp Gly Trp Ala Trp Pro Ser Val Gly Trp Gly Gly Gly Arg
 165 170 175

Ala Ser Ser Ala Arg Ser Ser Leu Arg Arg Gly Arg Ser Gln Gly Ser
 180 185 190

Gly Ala Arg Leu Val Gly Arg Glu Thr Pro Gly Ala Leu Ser Ser Gly
 195 200 205

Glu Val Gly Val Gln Ala Gly Lys Pro Gly Val Ser Arg Gly Ala Ala
 210 215 220

Val Arg Ser Arg Val Gln Gln Glu Gly Ser Pro Asp Gly Gln Val Pro
 225 230 235 240

Leu Ser Pro Gly Ala Gln His Trp Leu Val Ala Phe Ala Glu Val Val

245

250

255

Lys Lys Gly Arg Arg Pro Val Glu Arg Arg Ser Pro Gly Ile Pro Asn
 260 265 270

Tyr Val Thr Gly Asp Pro Phe Gln Phe Gly Leu Trp Pro Ile Pro Leu
 275 280 285

Ser Pro Pro Ala Asp Trp Ser Leu Leu Ser Gly Ser Pro Gln Pro Phe
 290 295 300

Leu Phe Asn Arg Gly Gln Arg Gly Asp Gly Glu Ser Thr Asp Gly Gly
 305 310 315 320

Cys Gly Ala Gly Glu Ala Ala Gly Arg Arg Ala Gly Leu Val Gly Arg
 325 330 335

Ala Gly Arg Val Gln Gly Phe Arg Val Thr Cys Pro Ala Pro Arg His
 340 345 350

Arg Ala Gly Arg Cys Ser Leu Pro Ile Cys Phe Arg Pro Ser Ser Arg
 355 360 365

Phe Arg Arg Arg Val Gly Thr Ala Glu Pro Arg Arg Ala Pro Gly Arg
 370 375 380

Arg Arg Gln Gly Ala Gly Ile Ala Leu Ala Ala Gly Arg Ala Ala Ser
 385 390 395 400

Pro Gly Glu Glu Arg Ser Arg Arg Arg Leu Pro Ser His Pro Ala Thr
 405 410 415

Pro Ala Ser Arg Arg Pro Phe Pro Arg Ser Ala Arg Gln Ile Gln Arg
 420 425 430

Leu Pro Gly Ala Gly Asp Gly Val Val Pro Thr Ala Glu Gly Trp Thr
 435 440 445

Leu Ser Met Ser Asp Ala Ala Cys Gly Gln Pro Tyr Pro Asn Pro Thr
 450 455 460

Ala His Pro Asp Asn Gln Asn Leu Val Arg Pro Pro Gly Ser Cys Leu
 465 470 475 480

Val Trp Ser Gln Val Leu Ser Ala Pro Ser Pro Gly Pro Phe Thr Leu
 485 490 495

Gln Glu Leu His Ala Pro Leu Thr Ser Ala Phe Pro Trp Gln Gln Arg
 500 505 510

Gly Phe Ala Gly Arg Pro Gly Ser Pro Glu His Ser Ser Pro Leu Pro
515 520 525

Gly Gly Leu Leu Ala Leu Ala Gly Asp Thr Ser Arg Ser Phe Lys Cys
530 535 540

Pro Leu Gln Ser Leu Ile Asn Asp Pro Ile His Ser Pro Leu Leu Ser
545 550 555 560

Phe Val Ser Ala Ile Glu Lys Cys Leu Pro Arg Ala Ala Leu His Phe
565 570 575

Arg Pro Leu Phe Cys Val Leu Leu
580